Page 1 of 8

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RAW SEQUENCE LISTING DATE: 11/23/2001 PATENT APPLICATION: US/09/899,303A TIME: 18:16:03

Input Set : N:\Crf3\RULE60\09899303A.raw Output Set: N:\CRF3\11232001\1899303A.raw

SEQUENCE LISTING

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ENTERED
      3 (1) GENERAL INFORMATION:
             (i) APPLICANT: MAERTENS, GEERT
                            BOSMAN, FONS
      7
                            DE MARTYNOFF, GUY
      8
                            BUYSE, MARIE-ANGE
     10
            (ii) TITLE OF INVENTION: PURIFIED HEPATITIS C VIRUS ENVELOPE
     11
                                     PROTEINS FOR DIAGNOSTIC AND THERAPEUTIC USE
     13
           (iii) NUMBER OF SEQUENCES: 111
     15
            (iv) CORRESPONDENCE ADDRESS:
     16
                  (A) ADDRESSEE: NIXON & VANDERHYE P.C.
     17
                  (B) STREET: 1100 NORTH GLEBE ROAD
     18
                  (C) CITY: ARLINGTON
     19
                  (D) STATE: VIRGINIA
     20
                  (E) COUNTRY: U.S.A.
     21
                  (F) ZIP: 22201-4714
     23
             (v) COMPUTER READABLE FORM:
     24
                  (A) MEDIUM TYPE: Floppy disk
     25
                  (B) COMPUTER: IBM PC compatible
                  (C) OPERATING SYSTEM: PC-DOS/MS-DOS
     26
     27
                  (D) SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
     29
            (vi) CURRENT APPLICATION DATA:
C-->, 30
                  (A) APPLICATION NUMBER: US/09/899,303A
C--> 31
                  (B) FILING DATE: 06-Jul-2001
     32
                  (C) CLASSIFICATION:
     34
           (vii) PRIOR APPLICATION DATA:
     35
                  (A) APPLICATION NUMBER: US/08/612,973
     36
                  (B) FILING DATE: 11-MAR-1996
     38
          (viii) ATTORNEY/AGENT INFORMATION:
     39
                  (A) NAME: BYRNE, THOMAS E.
     40
                  (B) REGISTRATION NUMBER: 32,205
     41
                  (C) REFERENCE/DOCKET NUMBER: 1487-10
     43
            (ix) TELECOMMUNICATION INFORMATION:
     44
                  (A) TELEPHONE: (703) 816-4000
     45
                  (B) TELEFAX: (703) 816-4100
     49 (2) INFORMATION FOR SEQ ID NO: 1:
             (i) SEQUENCE CHARACTERISTICS:
     52
                  (A) LENGTH: 21 base pairs
     53
                  (B) TYPE: nucleic acid
                  (C) STRANDEDNESS: single
     54
     55
                  (D) TOPOLOGY: linear
     57
            (ii) MOLECULE TYPE: cDNA
     59
           (iii) HYPOTHETICAL: NO
C--> 61
            (iv) ANTI-SENSE: NO
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
     68 GGCATGCAAG CTTAATTAAT T
                                                                                 21
     70 (2) INFORMATION FOR SEQ ID NO: 2:
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	72		(i)	SEQ	JENC	E CHA	ARACI	CERIS	STICS	3:								
	73			(A)) LEI	NGTH:	: 68	base	e pai	irs								
	74			(B	TY!	PE: r	nucle	eic a	acid						,			
	75			(C) ST	RANDI	EDNES	SS: 8	singl	Le								
	76				TOI													
	78	(ii)	MOLI	CUL	TYI	PE: 0	DNA										
	80				OTHE													
c>	82	•			I-SEN													
	86	,	•					OITS	N: SI	EO II	ON C	: 2:						
	88												CACC	ATCA	T A	ATAG:	TAAT	60
		TAACI																68
	92	(2) 1	NFO	RMAT	ION I	OR S	SEQ 1	D NO): 3:	:								
	94	` '																
	95	· · ·																
	96																	
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	100		(ii)) MOI	LECUI	LE TY	PE:	cDN	A									
	102	(POTHE													
C>	104	`	(iv	AN'	ri-si	ENSE:	NO:											
	107				ATURE													
	108																	
	109	109 (B) LOCATION: 1639																
	111 (ix) FEATURE:																	
	112			(2	A) NA	AME/I	KEY:	mat_	_pept	tide								
	113			(I	3) LO	CAT	ON:	16	36									
	116		(xi)) SE	QUENC	CE DI	ESCR	[PTIC	ON: S	SEQ :	ID NO): 3	:					
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	119	Met	${\tt Pro}$	Gly	Cys	Ser	Phe	Ser	Ile	Phe	Leu	Leu	Ala	Leu	Leu	Ser	Cys	
	120	1				5					10					15		
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	123	Leu	Thr	Ile	Pro	Ala	Ser	Ala	Tyr	Glu	Val	Arg	Asn	Val	Ser	Gly	Met	
	124				20					25					30			
	126	TAC	CAT	GTC	ACG	AAC	GAC	TGC	TCC	AAC	TCA	AGC	ATT	GTG	TAT	GAG	GCA	144
		Tyr	His	Val	Thr	Asn	Asp	Cys	Ser	Asn	Ser	Ser	Ile	Val	\mathtt{Tyr}	Glu	Ala	
	128			35					40					45				
		GCG																192
		Ala			Ile	Met			Pro	Gly	Cys			Cys	Val	Arg	Glu	
	132		50					55					60					
	134	AAC	AAC	TCT	TCC	CGC	TGC	TGG	GTA	GCG	CTC	ACC	CCC	ACG	CTC	GCA	GCT	240
		Asn	Asn	Ser	Ser	Arg	Cys	\mathtt{Trp}	Val	Ala	Leu	Thr	Pro	Thr	Leu	Ala	Ala	
		65					70					75					80	
	138	AGG	AAC	GCC	AGC	GTC	CCC	ACC	ACG	ACA	ATA	CGA	CGC	CAC	GTC	GAT	TTG	288
		Arg	Asn	Ala	Ser	Val	Pro	Thr	Thr	Thr		Arg	Arg	His	Val	_	Leu	
	140					85					90					95		
		CTC																336
		Leu	Val	Gly		Ala	Ala	Leu	Cys		Ala	Met	\mathtt{Tyr}	Val	Gly	Asp	Leu	
	144				100					105					110			
	146	TGC	GGA	TCT	GTC	TTC	CTC	GTC	TCC	CAG	CTG	TTC	ACC	ATC	TCG	CCT	CGC	384

	Cys	Gly		Val	Phe	Leu	Val		Gln	Leu	Phe	Thr		Ser	Pro	Arg		
148	000	C a m	115	N CC	CITIC	CAC	CAC	120	3 3 M	mcc.	шса	אחכ	125	ccc	cca	CAC		422
	CGG																	432
152	Arg	130	GIU	1111	Vai	GIII	135	Cys	ASII	Cys	ser	140	TYL	PIO	GIY	urs		
	a ma		CCM	CAC	CCT	አመረ		mcc	CAM	እ መረገ	a mc		A A C	mcc	mcc	CCM		480
	ATA Ile																	400
	145	1111	GIY	птэ	AIG	150	MIG	пр	кър	Met	155	Mec	ASII	пр	361	160		
	ACA	ACC.	GCC	СТС	стс		TCG	CAG	СТС	СТС		አ ጥር	CCA	CAA	CCT			528
	Thr																	320
160					165			· · · ·		170	5			· · · ·	175			
	GTG	GAC	ATG	GTG		GGG	GCC	CAT	TGG		GTC	CTG	GCG	GGC		GCC		576
	Val																	
164		•		180		-			185	•				190				
166	TAC	TAT	TCC	ATG	GTG	GGG	AAC	TGG	GCT	AAG	GTT	TTG	ATT	GTG	ATG	CTA		624
	Tyr																	
168	-	_	195			_		200		_			205					
170	CTC	TTT	GCT	CTC	TAA	ľAG												642
171	Leu	Phe	Ala	Leu														
172		210									•							
175	(2)	INF	ORMA	rion	FOR	SEQ	ID I	: ON	4:					•				
177		(i)) SE(QUEN	CE CI	HARA	CTER	ISTI	CS:	•								
178			(2	A) LI	ENGTI	H: 2	12 ar	nino	acio	ds	•							
179					YPE:													
180		•	(I) T	OPOL	OGY:	line	ear										
182) MOI															
184) SE															
	Met	Pro	Gly	Cys		Phe	Ser	Ile	Phe		Leu	Ala	Leu	Leu		Cys		
187	_ 1			_	. 5	_		_		10	_	_		_	15		•	
	Leu	Thr	IIe		Ala	Ser	Ala	Tyr		Val	Arg	Asn	vaı			Met		
190	m	774 -	17- 1	20	3	3	0	C	25	C	O	T1 -	37 <u>-</u> 3	30		230		
192	Tyr	HIS	35	THE	ASI	Asp	Cys	40	ASII	ser	ser	тте		туг	GIU	Ald		
	Ala	7 an	-	Tlo	Mo+	Wi c	Πh~		C111	CTTC	wa 1	Dro	45 Cvc	Wal.	λνα	Clu		
196	Ala	50	Met	116	Met	urs	55	PIO	GIY	Cys	vai	60	Cys	vaı	AIG	GIU		
	Asn		Ser	Ser	Δrσ	Cvs		Val	Δla	T.e.ii	Thr		Thr	T.e.u	Δla	Δla		
199			001	001	*** 9	70		, ,	····u	Leu	75	110	1111	ДСИ		80		
	Arg	Asn	Ala	Ser	Val		Thr	Thr	Thr	Ile		Arσ	His	Val	Asp			
202	_									90	_	5			95			
	Leu	Val	Glv	Ala								Tvr	Val	Glv		Leu		
205			1	100				-1-	105			- 4 -		110				
	Cys	Gly	Ser	Val	Phe	Leu	Val	Ser	Gln	Leu	Phe	Thr	Ile	Ser	Pro	Arg		
208		•	115	_				120	_	_		•	125			_		
	Arg	His		Thr	Val	Gln	Asp		Asn	Cys	Ser	Ile		Pro	Gly	His		
211		130					135			•		140	-		-			
213	Ile		Gly	His	Arg	Met	Ala	Trp	Asp	Met	Met		Asn	Trp	Ser	Pro		
	145		-			150		_	-		155			-		160		
216	Thr	Thr	Ala	Leu	Val	Val	Ser	Gln	Leu	Leu	Arg	Ile	Pro	Gln	Ala	Val		
217		•			165					170	_				175			

	219 220	Val	Asp	Met	Val 180	Ala	Gly	Ala	His	Trp 185	Gly	Val	Leu	Ala	Gly 190	Leu	Ala	
		Tyr	Tyr	Ser 195		Val	Gly	Asn	Trp 200	-	Lys	Val	Leu			Met	Leu	
		T.e.11	Phe	Ala	Len				200					205				
	226	Dea	210	nia	шец													
		(2)		ORMA	rion	FOR	SEQ	ID I	NO: 5	5 :								
	231	•							ISTIC									
	232			(]	A) LI	ENGT	H: 79	95 ba	ase p	pairs	5							
	233			(I	B) T	YPE:	nuc	leic	acio	ī								
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	248		(ix)		ATURI						,							
	249		•	(<i>I</i>	A) NZ	AME/I	KEY:	mat_	_pept	ide								
	250			· (I	3) LO	CAT:	ION:	1	789									
	253		(xi) SE(QUEN	CE DI	ESCR	IPTI	ON: S	SEQ :	D NO	D: 5	:					
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			Leu	Gly	Lys		Ile	Asp	Thr	Leu		Cys	Gly	Phe	Ala		Leu	
	257	1	~~~	m» a	3 000	5	ama	C.T.C	000	222	10	CEN.	000	000		15	200	0.6
									GGC Gly									96
	261	Val	Gry	тут	20	PIO	ьеи	val	GIY	25	PIO	ьeu	GIY	GTĀ	30	АТА	AIG	
		GCC	СТС	GCG		GGC	GTC	CGG	GTT		GAG	GAC	GGC	GTG		тат	GCA	144
									Val									
	265			35		•		_	40			-	-	45		-		
	267	ACA	GGG	AAT	TTG	CCC	GGT	TGC	TCT	TTC	TCT	ATC	TTC	CTC	TTG	GCT	TTG	192
		Thr	_	Asn	Leu	Pro	Gly	Cys	Ser	Phe,	Ser	Ile	Phe	Leu	Leu	Ala	Leu	
	269		50					55					60					
									GCT									240
			Ser	Cys	Leu	Thr		Pro	Ala	Ser	Ala	_	GIu	Val	Arg	Asn		
	273	65 TCC	ccc	አ ሞር	መአ 🔿	CNT	70	N.C.C	AAC	CAC	TIC C	75	7 7 C	mc x	700	א חיים	80	288
									Asn									200
	277	501		1100	-1-	85	vai	1111	non	nop	90	501	ASII	501	DCI	95	var	
		TAT	GAG	GCA	GCG		ATG	ATC	ATG	CAC		CCC	GGG	TGC	GTG		TGC	336
									Met									
	281				100					105			_	_	110		_	
									CGC									384
		Val	Arg		Asn	Asn	Ser	Ser	Arg	Cys	Trp	Val	Ala		Thr	Pro	Thr	
	285			115					120					125				
									GTC									432
		ьeu		АТА	Arg	Asn	Ala		Val	Pro	Thr	Thr		Ile	Arg	Arg	Hls	
	289		130					135					140					

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	145	a. a	cm.c	maa		150	с тс	mm-c	ama.		155	63.6	ama			160		
															ACC			528
	GIĀ	Asp	ьeu	Cys	-	Ser	val	Pne	Leu		Ser	GIn	Leu	Pne	Thr	ile		
297					165					170					175			
															ATC			576
	Ser	Pro	Arg	-	HIS	GLu	Thr	vaı		Asp	Cys	Asn	Cys		Ile	Tyr		
301	000		~~~	180			~ 3 ~		185		maa	a		190	3 550			
															ATG			624
	Pro	GTA		TTE	Thr	GTA	HIS		мет	Ата	Trp	Asp		мет	Met	Asn		
305	maa.	шаа	195	3.03			ama.	200	C III 3	maa	a a	ama	205	000	3 000			c70
															ATC			672
	Trp		Pro	THE	THE	Ата		val	vaı	ser	GIN		Leu	Arg	Ile	Pro		
309	CD 3	210	CITIC	CITIC	C3 C	3 mc	215	coc	CCC	600	(13 M	220	CCA	cma	СШС	cac	-	720
															CTG			720
		Ala	Val	val	Asp		val	Ald	GTA	Ald		тгр	GIY	Val	Leu			
	225	ama	000	ma a	mam	230	3 mc	cmc	ccc	330	235	COM	3 3 C	cmm	ттс	240		760
															TTG			768
317	GIY	ьeu	АІА	TYL	245	Ser	Met	val	GIY		ттр	Ald	гÃг	Vdl	Leu	TTE		•
	GTG	አ መረግ	CIDIX	CITIC		CCIII	ccc	חאאר	Πλ <i>C</i>	250					255			795
	_				_	_		TAA.	LAG									793
	Val	Met	Leu		Pne	Ald	Pro											
321	(2)	TNE	יאשמי	260	EVD	CEO.	TD 1	viO - 4	ε.									
324	(2)	TML	JRIM.	LION	LOK	350	10 1	NO: 0	3 .									
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326		(i		_		HARA	CTER	ISTIC	cs:	3.c								
327		(i	(1	A) LI	ENGTI	HARAG	CTER: 53 ar	ISTIC mino	cs:	is								
327 328		(i	(<i>I</i>	A) LI 3) T	ENGTI PE:	HARAC H: 26	CTER: 53 ar 10 ac	ISTIC mino cid	cs:	ls								
327 328 329		•	(A) LI B) TY O) T(ENGTI PE: OPOLO	HARAGH: 26 amir DGY:	CTER: 53 ar no ac line	ISTIO mino cid ear	cs:	is								
327 328 329 331		(ii	I) I) IOM (A) LI 3) TI 0) T(LECUI	ENGTI (PE: OPOLO LE TY	HARACH: 26 amir DGY:	CTER: 53 ar no ac line prof	ISTIC mino cid ear tein	CS: acid) · 6							
327 328 329 331 333		(ii (xi	(/ (I (I) MOI) SE(A) LI B) T C) T C LECUI QUEN	ENGTI (PE: OPOLO LE TI CE DI	HARACH: 26 amin DGY: YPE: ESCRI	CTER: 53 ar no ac line prof	ISTIC mino cid ear tein	CS: acid	ED NO			Phe	Ala	Asn	T.e.u		
327 328 329 331 333 335	Met	(ii (xi	(/ (I (I) MOI) SE(A) LI B) T C) T C LECUI QUEN	ENGTH (PE: OPOLO LE TY CE DI Val	HARACH: 26 amin DGY: YPE: ESCRI	CTER: 53 ar no ac line prof	ISTIC mino cid ear tein	CS: acid	ID NO Thr			Phe	Ala	Asp 15	Leu		
327 328 329 331 333 335 336	Met 1	(ii (xi Leu	(P (I (I) MOI) SE(Gly	A) LI B) TY C) T(LECUI QUEN(LYS	ENGTI (PE: OPOLO LE TY CE DI Val	HARAC H: 26 amin DGY: YPE: ESCRI	CTER: 53 ar no ac line prot IPTIC Asp	ISTIC mino cid ear tein DN: S	CS: acio SEQ : Leu	ID NO Thr 10	Cys	Gly			15			
327 328 329 331 333 335 336 338	Met 1	(ii (xi Leu	(P (I (I) MOI) SE(Gly	A) LECUI QUENC LYS	ENGTI (PE: OPOLO LE TY CE DI Val	HARAC H: 26 amin DGY: YPE: ESCRI	CTER: 53 ar no ac line prot IPTIC Asp	ISTIC mino cid ear tein DN: S	SEQ :	ID NO Thr 10	Cys	Gly		Ala	_			
327 328 329 331 333 335 336 338 339	Met 1 Val	(ii (xi Leu Gly	(I (I (I)) MOI) SEG Gly	A) LH B) TO COLUMN CUENC LYS Ile 20	ENGTH YPE: DPOLO LE TY CE DI Val 5 Pro	HARACH: 26 amin DGY: VPE: ESCRI	CTER: 53 ar no ac line prod IPTIC Asp Val	ISTIC mino cid ear tein DN: S Thr	SEQ : Leu Ala 25	ID NO Thr 10 Pro	Cys Leu	Gly Gly	Gly	Ala 30	15 Ala	Arg		
327 328 329 331 333 335 336 338 339	Met 1 Val	(ii (xi Leu Gly	(I (I (I)) MOI) SEG Gly	A) LH B) TO COLUMN CUENC LYS Ile 20	ENGTH YPE: DPOLO LE TY CE DI Val 5 Pro	HARACH: 26 amin DGY: VPE: ESCRI	CTER: 53 ar no ac line prod IPTIC Asp Val	ISTIC mino cid ear tein DN: S Thr	SEQ : Leu Ala 25	ID NO Thr 10 Pro	Cys Leu	Gly Gly	Gly	Ala 30	15	Arg		
327 328 329 331 333 335 336 338 341 342	Met 1 Val	(ii (xi Leu Gly	(1) (I) (I) (I) (I) (I) (I) (I) (I) (I) (I	A) LH B) TY C) TO LECUI QUENC Lys Ile 20 His	ENGTH YPE: DPOLO LE TY CE DH Val 5 Pro	HARAG H: 26 amin DGY: YPE: ESCRI Ile Leu Val	CTER: 53 ar 10 ac 1ine prof IPTIC Asp Val	ISTIC mino cid ear tein ON: S Thr Gly Val 40	SEQ : Leu Ala 25 Leu	ID NO Thr 10 Pro Glu	Cys Leu Asp	Gly Gly	Gly Val 45	Ala 30 Asn	15 Ala Tyr	Arg Ala		
327 328 329 331 333 335 336 338 341 342 344	Met 1 Val Ala Thr	(ii (xi Leu Gly Leu	(1) (I) (I) (I) (I) (I) (I) (I) (I) (I) (I	A) LH A) TH B) TO COMMENT COMMENT COMMENT LYS LYS LYS LH LE LO His	ENGTH YPE: DPOLO LE TY CE DH Val 5 Pro	HARACHE 26 amin DGY: YPE: ESCRITLE Leu Val	CTER: 53 ar 10 ac 1ine prof IPTIC Asp Val Arg	ISTIC mino cid ear tein ON: S Thr Gly Val 40	SEQ Theu Ala 25 Leu Phe	ID NO Thr 10 Pro Glu Ser	Cys Leu Asp Ile	Gly Gly Gly Phe	Gly Val 45 Leu	Ala 30 Asn	15 Ala	Arg Ala		
327 328 329 331 333 335 336 338 341 342 344 345	Met 1 Val Ala Thr	(ii (xi Leu Gly Leu Gly 50	(I (I (I)) MOI) SEG Gly Tyr Ala 35 Asn	A) LH B) TY D) TO LECUI QUENO Lys Ile 20 His	ENGTH YPE: DPOLO LE TY CE DI Val 5 Pro Gly	HARACH: 26 amin DGY: YPE: ESCRI Ile Leu Val Gly	CTER: 53 ar 10 ac 1ine prod IPTIC Asp Val Arg Cys 55	ISTIC mino cid ear tein ON: S Thr Gly Val 40 Ser	SEQ : Leu Ala 25 Leu Phe	ID NO Thr 10 Pro Glu Ser	Cys Leu Asp Ile	Gly Gly Gly Phe 60	Gly Val 45 Leu	Ala 30 Asn Leu	15 Ala Tyr Ala	Arg Ala Leu		
327 328 329 331 333 335 336 338 341 342 344 345	Met 1 Val Ala Thr	(ii (xi Leu Gly Leu Gly 50	(I (I (I)) MOI) SEG Gly Tyr Ala 35 Asn	A) LH B) TY D) TO LECUI QUENO Lys Ile 20 His	ENGTH YPE: DPOLO LE TY CE DI Val 5 Pro Gly	HARACH: 26 amin DGY: YPE: ESCRI Ile Leu Val Gly	CTER: 53 ar 10 ac 1ine prod IPTIC Asp Val Arg Cys 55	ISTIC mino cid ear tein ON: S Thr Gly Val 40 Ser	SEQ : Leu Ala 25 Leu Phe	ID NO Thr 10 Pro Glu Ser	Cys Leu Asp Ile	Gly Gly Gly Phe 60	Gly Val 45 Leu	Ala 30 Asn Leu	15 Ala Tyr	Arg Ala Leu		
327 328 329 331 333 335 336 338 341 342 344 345 347 348	Met 1 Val Ala Thr Leu 65	(ii (xi Leu Gly Leu Gly 50 Ser	(I (I (I) MOI) SEG Gly Tyr Ala 35 Asn	A) LH B) TY C) TC CUIECUI LYS LYS 11e 20 His Leu	ENGTH YPE: DPOLO LE TY CE DH Val 5 Pro Gly Pro Thr	HARACH: 26 amin DGY: VPE: ESCRI ILe Leu Val Gly Val 70	CTER: 53 ar 10 ac 1ine prof PTIC Asp Val Arg Cys 55 Pro	ISTIC mino cid ear tein ON: S Thr Gly Val 40 Ser	SEQ : Leu Ala 25 Leu Phe Ser	ID No Thr 10 Pro Glu Ser Ala	Cys Leu Asp Ile Tyr 75	Gly Gly Gly Phe 60 Glu	Gly Val 45 Leu Val	Ala 30 Asn Leu Arg	15 Ala Tyr Ala Asn	Arg Ala Leu Val 80		
327 328 329 331 333 335 336 338 341 342 344 345 347 348	Met 1 Val Ala Thr Leu 65	(ii (xi Leu Gly Leu Gly 50 Ser	(I (I (I) MOI) SEG Gly Tyr Ala 35 Asn	A) LH B) TY C) TC CUIECUI LYS LYS 11e 20 His Leu	ENGTH YPE: DPOLO LE TY CE DH Val 5 Pro Gly Pro Thr	HARACH: 26 amin DGY: VPE: ESCRI ILe Leu Val Gly Val 70	CTER: 53 ar 10 ac 1ine prof PTIC Asp Val Arg Cys 55 Pro	ISTIC mino cid ear tein ON: S Thr Gly Val 40 Ser	SEQ : Leu Ala 25 Leu Phe Ser	ID No Thr 10 Pro Glu Ser Ala	Cys Leu Asp Ile Tyr 75	Gly Gly Gly Phe 60 Glu	Gly Val 45 Leu Val	Ala 30 Asn Leu Arg	15 Ala Tyr Ala	Arg Ala Leu Val 80		
327 328 329 331 333 335 336 338 341 342 344 345 347 348 350 351	Met 1 Val Ala Thr Leu 65 Ser	(ii (xi Leu Gly Leu Gly 50 Ser	(I (I (I)) MOI) SEG Gly Tyr Ala 35 Asn Cys	A) LH B) TY C) TO LECUI QUENC Lys Ile 20 His Leu Leu Tyr	ENGTH YPE: PPOLO LE TY Val 5 Pro Gly Pro Thr His 85	HARACH: 26 amin DGY: VPE: ESCR: Ile Leu Val Gly Val 70 Val	CTER: 53 ar 53 ar 10 ac 11 ne prof 1PTIC Asp Val Arg Cys 55 Pro Thr	ISTIC mino cid ear tein ON: S Thr Gly Val 40 Ser Ala	SEQ : Leu Ala 25 Leu Phe Ser	ID No Thr 10 Pro Glu Ser Ala Cys 90	Cys Leu Asp Ile Tyr 75 Ser	Gly Gly Gly Phe 60 Glu Asn	Gly Val 45 Leu Val Ser	Ala 30 Asn Leu Arg	15 Ala Tyr Ala Asn Ile 95	Arg Ala Leu Val 80 Val		
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327 328 329 331 333 335 336 338 341 342 344 345 347 348 350 351 353 354 356 357	Met 1 Val Ala Thr Leu 65 Ser Tyr	(ii (xi Leu Gly 50 Ser Gly Glu Arg	(I (I (I) MOI) SE(Gly Tyr Ala 35 Asn Cys Met Ala Glu 115	A) LH B) TY C) TC CUI CUENC Lys Ile 20 His Leu Leu Tyr Ala 100 Asn	ENGTH YPE: PPOLO CE DI Val Pro Gly Pro Thr His 85 Asp	HARACH: 26 amin DGY: VPE: ESCRI Ile Leu Val Gly Val 70 Val Met Ser	CTER: 53 ar 10 ac 1ine prod IPTIC Asp Val Arg Cys 55 Pro Thr Ile Ser	ISTIC mino cid ear tein ON: S Thr Gly Val 40 ser Ala Asn Met Arg 120	SEQ 1 Leu Ala 25 Leu Phe Ser Asp His 105 Cys	ID NO Thr 10 Pro Glu Ser Ala Cys 90 Thr	Cys Leu Asp Ile Tyr 75 Ser Pro Val	Gly Gly Phe 60 Glu Asn Gly Ala	Gly Val 45 Leu Val Ser Cys Leu 125	Ala 30 Asn Leu Arg Ser Val 110 Thr	15 Ala Tyr Ala Asn Ile 95 Pro	Arg Ala Leu Val 80 Val Cys		

VERIFICATION SUMMARY DATE: 11/23/2001 PATENT APPLICATION: US/09/899,303A TIME: 18:16:04

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